1. Title: US-10-615-383A-7_COPY_102_2894
Sequence: 1 ttaaaaaaaaataatttact.....atagaaaaaataaaaattaa 2793
DE Staphylococcus epidermidis ORF nucleic acid sequence SEQ ID NO:2477.
XX
KW Staphylococcus epidermidis; open reading frame; ORF; bacterial infection;
KW antibacterial; gene therapy; gene; ds.
XX
OS Staphylococcus epidermidis.
XX
PN US6380370-B1.
XX
PD 30-APR-2002.
XX
PF 13-AUG-1998; 98US-00134001.
XX
PR 14-AUG-1997; 97US-0055779P.
PR 08-NOV-1997; 97US-0064964P.
XX
PA (GENO-) GENOME THERAPEUTICS CORP.
XX
PI Doucette-Stamm LA, Bush D;
XX
DR WPI; 2002-381255/41.
DR P-PSDB; ABP40469.
XX
PT Novel isolated nucleic acid encoding a Staphylococcus epidermis
PT polypeptide, useful for diagnosing and treating bacterial infections.
XX
PS Disclosure; SEQ ID NO 2477; 267pp; English.
XX
CC ABN90538 to ABN93374 represent Staphylococcus epidermidis open reading
CC frame (ORF) nucleic acid sequences which encode the amino acid sequences
CC given in ABP35124 to ABP37960. The S. epidermidis sequences have
CC antibacterial activity and can be used in gene therapy. The sequences can
CC also be used in the diagnosis and treatment of bacterial infections,
CC particularly S. epidermidis infections. The sequences can be used to
CC screen for compounds able to interfere with the S. epidermidis life cycle
CC or inhibit S. epidermidis infection. N.B. The sequence data for this
CC patent did not form part of the printed specification, but was obtained
CC in electronic format directly from the USPTO web site
XX
SQ Sequence 2793 BP; 1149 A; 423 C; 497 G; 724 T; 0 U; 0 Other;

Query Match 99.9%; Score 2791.4; DB 1; Length 2793;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 2792; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy          1 TTAAAAAAAAAATAATTTACTAACTAAAAAGAAACCTATAGCAAATAAATCCAATAAATAT 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 TTAAAAAAAAAATAATTTACTAACTAAAAAGAAACCTATAGCAAATAAATCCAATAAATAT 60

Qy         61 GCAATTAGAAAATTCACAGTAGGTACAGCGTCTATTGTAATAGGTGCAGCATTATTGTTT 120
              |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db         61 GCAATTAGAAAATTCACAGTAGGTACAGCGTCTATTGTAATAGGTGCAACATTATTGTTT 120

Qy        121 GGTTTAGGTCATAATGAGGCCAAAGCTGAGGAGAATACAGTACAAGACGTTAAAGATTCG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 GGTTTAGGTCATAATGAGGCCAAAGCTGAGGAGAATACAGTACAAGACGTTAAAGATTCG 180

Qy        181 AATATGGATGATGAATTATCAGATAGCAATGATCAGTCCAGTAATGAAGAAAAGAATGAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 AATATGGATGATGAATTATCAGATAGCAATGATCAGTCCAGTAATGAAGAAAAGAATGAT 240
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Qy       241 GTAATCAATAATAGTCAGTCAATAAACACCGATGATGATAACCAAATAAAAAAAGAAGAA 300
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       241 GTAATCAATAATAGTCAGTCAATAAACACCGATGATGATAACCAAATAAAAAAAGAAGAA 300

Qy       301 ACGAATAGCAACGATGCCATAGAAAATCGCTCTAAAGATATAACACAGTCAACAACAAAT 360
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       301 ACGAATAGCAACGATGCCATAGAAAATCGCTCTAAAGATATAACACAGTCAACAACAAAT 360

Qy       361 GTAGATGAAAACGAAGCAACATTTTTTACAAAAGACCCCTCAAGATAATACTCAGCTTAAA 420
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       361 GTAGATGAAAACGAAGCAACATTTTTTACAAAAGACCCCTCAAGATAATACTCAGCTTAAA 420

Qy       421 GAAGAAGTGGTAAAAGAACCCTCATCAGTCGAATCCTCAAATTCATCAATGGATACTGCC 480
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       421 GAAGAAGTGGTAAAAGAACCCTCATCAGTCGAATCCTCAAATTCATCAATGGATACTGCC 480

Qy       481 CAACAACCATCTCATACAACAATAAATAGTGAAGCATCTATTCAAACAAGTGATAATGAA 540
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       481 CAACAACCATCTCATACAACAATAAATAGTGAAGCATCTATTCAAACAAGTGATAATGAA 540

Qy       541 GAAAATTCCCGCGTATCAGATTTTGCTAACTCTAAAATAATAGAGAGTAACACTGAATCC 600
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       541 GAAAATTCCCGCGTATCAGATTTTGCTAACTCTAAAATAATAGAGAGTAACACTGAATCC 600

Qy       601 AATAAAGAAGAGAATACTATAGAGCAACCTAACAAAGTAAGAGAAGATTCAATAACAAGT 660
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       601 AATAAAGAAGAGAATACTATAGAGCAACCTAACAAAGTAAGAGAAGATTCAATAACAAGT 660

Qy       661 CAACCGTCTAGCTATAAAAATATAGATGAAAAAATTTCAAATCAAGATGAGTTATTAAAT 720
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       661 CAACCGTCTAGCTATAAAAATATAGATGAAAAAATTTCAAATCAAGATGAGTTATTAAAT 720

Qy       721 TTACCAATAAATGAATATGAAAATAAGGTTAGACCGTTATCTACAACATCTGCCCAACCA 780
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       721 TTACCAATAAATGAATATGAAAATAAGGTTAGACCGTTATCTACAACATCTGCCCAACCA 780

Qy       781 TCGAGTAAGCGTGTAACCGTAAATCAATTAGCGGCAGAACAAGGTTCGAATGTTAATCAT 840
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       781 TCGAGTAAGCGTGTAACCGTAAATCAATTAGCGGCAGAACAAGGTTCGAATGTTAATCAT 840

Qy       841 TTAATTAAAGTTACTGATCAAAGTATTACTGAAGGATATGATGATAGTGATGGTATTATT 900
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       841 TTAATTAAAGTTACTGATCAAAGTATTACTGAAGGATATGATGATAGTGATGGTATTATT 900

Qy       901 AAAGCACATGATGCTGAAAACTTAATCTATGATGTAACTTTTGAAGTAGATGATAAGGTG 960
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       901 AAAGCACATGATGCTGAAAACTTAATCTATGATGTAACTTTTGAAGTAGATGATAAGGTG 960

Qy       961 AAATCTGGTGATACGATGACAGTGAATATAGATAAGAATACAGTTCCATCAGATTTAACC 1020
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       961 AAATCTGGTGATACGATGACAGTGAATATAGATAAGAATACAGTTCCATCAGATTTAACC 1020

Qy      1021 GATAGTTTTGCAATACCAAAAATAAAAGATAATTCTGGAGAAATCATCGCTACAGGTACT 1080
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1021 GATAGTTTTGCAATACCAAAAATAAAAGATAATTCTGGAGAAATCATCGCTACAGGTACT 1080

Qy      1081 TATGACAACACAAATAAACAAATTACCTACACTTTTACAGATTATGTAGATAAATATGAA 1140
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1081 TATGACAACACAAATAAACAAATTACCTACACTTTTACAGATTATGTAGATAAATATGAA 1140

Qy      1141 AATATTAAAGCGCACCTTAAATTAACATCATACATTGATAAATCAAAGGTTCCAAATAAT 1200
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1141 AATATTAAAGCGCACCTTAAATTAACATCATACATTGATAAATCAAAGGTTCCAAATAAT 1200
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2. Title:          US-10-615-383A-10_COPY_51_598
Sequence:        1 ENTVQDVKDSNMDDELSDSN.........TIAFSTSSGQGQGDLPPEKT 548
DE   Staphylococcus epidermidis ORF amino acid sequence SEQ ID NO:5314.
XX
KW   Staphylococcus epidermidis; open reading frame; ORF; bacterial infection;
KW   antibacterial; gene therapy.
XX
OS   Staphylococcus epidermidis.
XX
PN   US6380370-B1.
XX
PD   30-APR-2002.
XX
PF   13-AUG-1998;    98US-00134001.
XX
PR   14-AUG-1997;    97US-0055779P.
PR   08-NOV-1997;    97US-0064964P.
XX
PA   (GENO-) GENOME THERAPEUTICS CORP.
XX
PI   Doucette-Stamm LA, Bush D;
XX
DR   WPI; 2002-381255/41.
DR   N-PSDB; ABN93014.
XX
PT   Novel isolated nucleic acid encoding a Staphylococcus epidermis
PT   polypeptide, useful for diagnosing and treating bacterial infections.
XX
PS   Disclosure; SEQ ID NO 5314; 267pp; English.
XX
CC   ABN90538 to ABN93374 represent Staphylococcus epidermidis open reading
CC   frame (ORF) nucleic acid sequences which encode the amino acid sequences
CC   given in ABP35124 to ABP37960. The S. epidermidis sequences have
CC   antibacterial activity and can be used in gene therapy. The sequences can
CC   also be used in the diagnosis and treatment of bacterial infections,
CC   particularly S. epidermidis infections. The sequences can be used to
CC   screen for compounds able to interfere with the S. epidermidis life cycle
CC   or inhibit S. epidermidis infection. N.B. The sequence data for this
CC   patent did not form part of the printed specification, but was obtained
CC   in electronic format directly from the USPTO web site
XX
SQ   Sequence 930 AA;

  Query Match             100.0%;  Score 2808;  DB 1;  Length 930;
  Best Local Similarity   100.0%;  Pred. No. 2.3e-138;
  Matches  548;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy            1 ENTVQDVKDSNMDDELSDSNDQSSNEEKNDVINNSQSINTDDDNQIKKEETNSNDAIENR 60
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db           51 ENTVQDVKDSNMDDELSDSNDQSSNEEKNDVINNSQSINTDDDNQIKKEETNSNDAIENR 110

Qy           61 SKDITQSTTNVDENEATFLQKTPQDNTQLKEEVVKEPSSVESSNSSMDTAQQPSHTTINS 120
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          111 SKDITQSTTNVDENEATFLQKTPQDNTQLKEEVVKEPSSVESSNSSMDTAQQPSHTTINS 170

Qy          121 EASIQTSDNEENSRVSDFANSKIIESNTESNKEENTIEQPNKVREDSITSQPSSYKNIDE 180
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          171 EASIQTSDNEENSRVSDFANSKIIESNTESNKEENTIEQPNKVREDSITSQPSSYKNIDE 230

Qy          181 KISNQDELLNLPINEYENKVRPLSTTSAQPSSKRVTVNQLAAEQGSNVNHLIKVTDQSIT 240
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          231 KISNQDELLNLPINEYENKVRPLSTTSAQPSSKRVTVNQLAAEQGSNVNHLIKVTDQSIT 290
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Qy        241 EGYDDSDGIIKAHDAENLIYDVTFEVDDKVKSGDTMTVNIDKNTVPSDLTDSFAIPKIKD 300
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        291 EGYDDSDGIIKAHDAENLIYDVTFEVDDKVKSGDTMTVNIDKNTVPSDLTDSFAIPKIKD 350

Qy        301 NSGEIIATGTYDNTNKQITYTFTDYVDKYENIKAHLKLTSYIDKSKVPNNNTKLDVEYKT 360
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        351 NSGEIIATGTYDNTNKQITYTFTDYVDKYENIKAHLKLTSYIDKSKVPNNNTKLDVEYKT 410

Qy        361 ALSSVNKTITVEYQKPNENRTANLQSMFTNIDTKNHTVEQTIYINPLRYSAKETNVNISG 420
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        411 ALSSVNKTITVEYQKPNENRTANLQSMFTNIDTKNHTVEQTIYINPLRYSAKETNVNISG 470

Qy        421 NGDEGSTIIDDSTIIKVYKVGDNQNLPDSNRIYDYSEYEDVTNDDYAQLGNNNDVNINFG 480
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        471 NGDEGSTIIDDSTIIKVYKVGDNQNLPDSNRIYDYSEYEDVTNDDYAQLGNNNDVNINFG 530

Qy        481 NIDSPYIIKVISKYDPNKDDYTTIQQTVTMQTTINEYTGEFRTASYDNTIAFSTSSGQGQ 540
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        531 NIDSPYIIKVISKYDPNKDDYTTIQQTVTMQTTINEYTGEFRTASYDNTIAFSTSSGQGQ 590

Qy        541 GDLPPEKT 548
              ||||||||
Db        591 GDLPPEKT 598


Qy       1201 AACACTAAGTTAGATGTAGAATATAAGACGGCCCTTTCATCAGTAAATAAAACAATTACG 1260
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1201 AACACTAAGTTAGATGTAGAATATAAGACGGCCCTTTCATCAGTAAATAAAACAATTACG 1260

Qy       1261 GTTGAATATCAAAAACCTAACGAAAATCGGACTGCTAACCTTCAAAGTATGTTCACAAAC 1320
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1261 GTTGAATATCAAAAACCTAACGAAAATCGGACTGCTAACCTTCAAAGTATGTTCACAAAC 1320

Qy       1321 ATAGATACGAAAAACCATACAGTTGAGCAAACGATTTATATTAACCCTCTTCGTTATTCA 1380
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1321 ATAGATACGAAAAACCATACAGTTGAGCAAACGATTTATATTAACCCTCTTCGTTATTCA 1380

Qy       1381 GCCAAAGAAACAAATGTAAATATTTCAGGGAATGGCGATGAAGGTTCAACAATTATCGAC 1440
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1381 GCCAAAGAAACAAATGTAAATATTTCAGGGAATGGCGATGAAGGTTCAACAATTATCGAC 1440

Qy       1441 GATAGTACAATCATTAAAGTTTATAAGGTTGGAGATAATCAAAATTTACCAGATAGTAAC 1500
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1441 GATAGTACAATCATTAAAGTTTATAAGGTTGGAGATAATCAAAATTTACCAGATAGTAAC 1500

Qy       1501 AGAATTTATGATTACAGTGAATATGAAGATGTCACAAATGATGATTATGCCCAATTAGGA 1560
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1501 AGAATTTATGATTACAGTGAATATGAAGATGTCACAAATGATGATTATGCCCAATTAGGA 1560

Qy       1561 AATAATAATGACGTGAATATTAATTTTGGTAATATAGATTCACCATATATTATTAAAGTT 1620
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1561 AATAATAATGACGTGAATATTAATTTTGGTAATATAGATTCACCATATATTATTAAAGTT 1620

Qy       1621 ATTAGTAAATATGACCCTAATAAGGACGATTACACGACGATACAGCAAACTGTGACAATG 1680
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1621 ATTAGTAAATATGACCCTAATAAGGACGATTACACGACGATACAGCAAACTGTGACAATG 1680

Qy       1681 CAAACGACTATAAATGAGTATACTGGTGAGTTTAGAACAGCATCCTATGATAATACAATT 1740
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1681 CAAACGACTATAAATGAGTATACTGGTGAGTTTAGAACAGCATCCTATGATAATACAATT 1740

Qy       1741 GCTTTCTCTACAAGTTCAGGTCAAGGACAAGGTGACTTGCCTCCTGAAAAAACTTATAAA 1800
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Db       1741 GCTTTCTCTACAAGTTCAGGTCAAGGACAAGGTGACTTGCCTCCTGAAAAAACTTATAAA 1800
Qy       1801 ATCGGAGATTACGTATGGGAAGATGTAGATAAAGATGGTATTCAAAATACAAATGATAAT 1860
Db       1801 ATCGGAGATTACGTATGGGAAGATGTAGATAAAGATGGTATTCAAAATACAAATGATAAT 1860
Qy       1861 GAAAAACCGCTTAGTAATGTATTGGTAACTTTGACGTATCCTGATGGAACTTCAAAATCA 1920
Db       1861 GAAAAACCGCTTAGTAATGTATTGGTAACTTTGACGTATCCTGATGGAACTTCAAAATCA 1920
Qy       1921 GTCAGAACAGATGAAGAGGGGAAATATCAATTTGATGGGTTAAAAAACGGATTGACTTAT 1980
Db       1921 GTCAGAACAGATGAAGAGGGGAAATATCAATTTGATGGGTTAAAAAACGGATTGACTTAT 1980
Qy       1981 AAAATTACATTCGAAACACCGGAAGGATATACGCCGACGCTTAAACATTCAGGAACAAAT 2040
Db       1981 AAAATTACATTCGAAACACCGGAAGGATATACGCCGACGCTTAAACATTCAGGAACAAAT 2040
Qy       2041 CCTGCACTAGACTCAGAAGGCAATTCTGTATGGGTAACTATTAACGGACAAGACGATATG 2100
Db       2041 CCTGCACTAGACTCAGAAGGCAATTCTGTATGGGTAACTATTAACGGACAAGACGATATG 2100
Qy       2101 ACTATTGATAGCGGATTTTATCAAACACCTAAATATAGCTTAGGGAACTATGTATGGTAT 2160
Db       2101 ACTATTGATAGCGGATTTTATCAAACACCTAAATATAGCTTAGGGAACTATGTATGGTAT 2160
Qy       2161 GACACTAATAAAGATGGTATTCAAGGTGATGATGAAAAAGGAATCTCTGGAGTAAAAGTG 2220
Db       2161 GACACTAATAAAGATGGTATTCAAGGTGATGATGAAAAAGGAATCTCTGGAGTAAAAGTG 2220
Qy       2221 ACGTTAAAAGATGAAAACGGAAATATCATTAGTACAACAACAACTGATGAAAATGGAAAG 2280
Db       2221 ACGTTAAAAGATGAAAACGGAAATATCATTAGTACAACAACAACTGATGAAAATGGAAAG 2280
Qy       2281 TATCAATTTGATAATTTAAATAGTGGTAATTATATTGTTCATTTTGATAAACCTTCAGGT 2340
Db       2281 TATCAATTTGATAATTTAAATAGTGGTAATTATATTGTTCATTTTGATAAACCTTCAGGT 2340
Qy       2341 ATGACTCAAACAACAACAGATTCTGGTGATGATGACGAACAGGATGCTGATGGGGAAGAA 2400
Db       2341 ATGACTCAAACAACAACAGATTCTGGTGATGATGACGAACAGGATGCTGATGGGGAAGAA 2400
Qy       2401 GTCCATGTAACAATTACTGATCATGATGACTTTAGTATAGATAACGGATACTATGATGAC 2460
Db       2401 GTCCATGTAACAATTACTGATCATGATGACTTTAGTATAGATAACGGATACTATGATGAC 2460
Qy       2461 GACTCAGATTCAGATAGTGATTCAGACTCAGATAGCGACGACTCAGACTCCGATAGCGAT 2520
Db       2461 GACTCAGATTCAGATAGTGATTCAGACTCAGATAGCGACGACTCAGACTCCGATAGCGAT 2520
Qy       2521 TCCGACTCAGACAGCGACTCAGATTCCGATAGTGATTCAGATTCAGACAGTGACTCAGAC 2580
Db       2521 TCCGACTCAGACAGCGACTCAGATTCCGATAGTGATTCAGATTCAGACAGTGACTCAGAC 2580
Qy       2581 TCAGATAGTGATTCAGATTCAGACAGCGATTCCGACTCAGACAGTGACTCAGGATTAGAC 2640
Db       2581 TCAGATAGTGATTCAGATTCAGACAGCGATTCCGACTCAGACAGTGACTCAGGATTAGAC 2640
Qy       2641 AATAGCTCAGATAAGAATACAAAAGATAAATTACCGGATACAGGAGCTAATGAAGATCAT 2700
Db       2641 AATAGCTCAGATAAGAATACAAAAGATAAATTACCGGATACAGGAGCTAATGAAGATCAT 2700
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Qy       2701 GATTCTAAAGGCACATTACTTGGAGCTTTATTTGCAGGTTTAGGAGCGTTATTATTAGGG 2760
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2701 GATTCTAAAGGCACATTACTTGGAGCTTTATTTGCAGGTTTAGGAGCGTTATTATTAGGG 2760

Qy       2761 AAGCGTCGCAAAAATAGAAAAAATAAAAATTAA 2793
             |||||||||||||||||||||||||||||||||
Db       2761 AAGCGTCGCAAAAATAGAAAAAATAAAAATTAA 2793

3.  Title:           US-10-615-383A-7_COPY_252_1895
Sequence:        1 gagaatacagtacaagacgt.........acttgcctcctgaaaaaact  1644
DE    Staphylococcus epidermidis ORF nucleic acid sequence SEQ ID NO:2477.
XX
KW    Staphylococcus epidermidis; open reading frame; ORF; bacterial infection;
KW    antibacterial; gene therapy; gene; ds.
XX
OS    Staphylococcus epidermidis.
XX
PN    US6380370-B1.
XX
PD    30-APR-2002.
XX
PF    13-AUG-1998;    98US-00134001.
XX
PR    14-AUG-1997;    97US-0055779P.
PR    08-NOV-1997;    97US-0064964P.
XX
PA    (GENO-) GENOME THERAPEUTICS CORP.
XX
PI    Doucette-Stamm LA,   Bush D;
XX
DR    WPI; 2002-381255/41.
DR    P-PSDB; ABP40469.
XX
PT    Novel isolated nucleic acid encoding a Staphylococcus epidermis
PT    polypeptide, useful for diagnosing and treating bacterial infections.
XX
PS    Disclosure; SEQ ID NO 2477; 267pp; English.
XX
CC    ABN90538 to ABN93374 represent Staphylococcus epidermidis open reading
CC    frame (ORF) nucleic acid sequences which encode the amino acid sequences
CC    given in ABP35124 to ABP37960. The S. epidermidis sequences have
CC    antibacterial activity and can be used in gene therapy. The sequences can
CC    also be used in the diagnosis and treatment of bacterial infections,
CC    particularly S. epidermidis infections. The sequences can be used to
CC    screen for compounds able to interfere with the S. epidermidis life cycle
CC    or inhibit S. epidermidis infection. N.B. The sequence data for this
CC    patent did not form part of the printed specification, but was obtained
CC    in electronic format directly from the USPTO web site
XX
SQ    Sequence 2793 BP; 1149 A; 423 C; 497 G; 724 T; 0 U; 0 Other;

  Query Match             100.0%;  Score 1644;  DB 1;  Length 2793;
  Best Local Similarity   100.0%;  Pred. No. 3.3e-288;
  Matches 1644;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGAATACAGTACAAGACGTTAAAGATTCGAATATGGATGATGAATTATCAGATAGCAAT 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        151 GAGAATACAGTACAAGACGTTAAAGATTCGAATATGGATGATGAATTATCAGATAGCAAT 210

Qy         61 GATCAGTCCAGTAATGAAGAAAAGAATGATGTAATCAATAATAGTCAGTCAATAAACACC 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        211 GATCAGTCCAGTAATGAAGAAAAGAATGATGTAATCAATAATAGTCAGTCAATAAACACC 270
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Qy      121 GATGATGATAACCAAATAAAAAAAGAAGAAACGAATAGCAACGATGCCATAGAAAATCGC 180
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      271 GATGATGATAACCAAATAAAAAAAGAAGAAACGAATAGCAACGATGCCATAGAAAATCGC 330

Qy      181 TCTAAAGATATAACACAGTCAACAACAAATGTAGATGAAAACGAAGCAACATTTTTTACAA 240
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      331 TCTAAAGATATAACACAGTCAACAACAAATGTAGATGAAAACGAAGCAACATTTTTTACAA 390

Qy      241 AAGACCCCTCAAGATAATACTCAGCTTAAAGAAGAAGTGGTAAAAGAACCCTCATCAGTC 300
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      391 AAGACCCCTCAAGATAATACTCAGCTTAAAGAAGAAGTGGTAAAAGAACCCTCATCAGTC 450

Qy      301 GAATCCTCAAATTCATCAATGGATACTGCCCAACAACCATCTCATACAACAATAAATAGT 360
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      451 GAATCCTCAAATTCATCAATGGATACTGCCCAACAACCATCTCATACAACAATAAATAGT 510

Qy      361 GAAGCATCTATTCAAACAAGTGATAATGAAGAAAATTCCCGCGTATCAGATTTTGCTAAC 420
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      511 GAAGCATCTATTCAAACAAGTGATAATGAAGAAAATTCCCGCGTATCAGATTTTGCTAAC 570

Qy      421 TCTAAAATAATAGAGAGTAACACTGAATCCAATAAAGAAGAGAATACTATAGAGCAACCT 480
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      571 TCTAAAATAATAGAGAGTAACACTGAATCCAATAAAGAAGAGAATACTATAGAGCAACCT 630

Qy      481 AACAAAGTAAGAGAAGATTCAATAACAAGTCAACCGTCTAGCTATAAAAATATAGATGAA 540
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      631 AACAAAGTAAGAGAAGATTCAATAACAAGTCAACCGTCTAGCTATAAAAATATAGATGAA 690

Qy      541 AAAATTTCAAATCAAGATGAGTTATTAAATTTACCAATAAATGAATATGAAAATAAGGTT 600
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      691 AAAATTTCAAATCAAGATGAGTTATTAAATTTACCAATAAATGAATATGAAAATAAGGTT 750

Qy      601 AGACCGTTATCTACAACATCTGCCCAACCATCGAGTAAGCGTGTAACCGTAAATCAATTA 660
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      751 AGACCGTTATCTACAACATCTGCCCAACCATCGAGTAAGCGTGTAACCGTAAATCAATTA 810

Qy      661 GCGGCAGAACAAGGTTCGAATGTTAATCATTTAATTAAAGTTACTGATCAAAGTATTACT 720
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      811 GCGGCAGAACAAGGTTCGAATGTTAATCATTTAATTAAAGTTACTGATCAAAGTATTACT 870

Qy      721 GAAGGATATGATGATAGTGATGGTATTATTAAAGCACATGATGCTGAAAACTTAATCTAT 780
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      871 GAAGGATATGATGATAGTGATGGTATTATTAAAGCACATGATGCTGAAAACTTAATCTAT 930

Qy      781 GATGTAACTTTTGAAGTAGATGATAAGGTGAAATCTGGTGATACGATGACAGTGAATATA 840
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      931 GATGTAACTTTTGAAGTAGATGATAAGGTGAAATCTGGTGATACGATGACAGTGAATATA 990

Qy      841 GATAAGAATACAGTTCCATCAGATTTAACCGATAGTTTTGCAATACCAAAAATAAAAGAT 900
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      991 GATAAGAATACAGTTCCATCAGATTTAACCGATAGTTTTGCAATACCAAAAATAAAAGAT 1050

Qy      901 AATTCTGGAGAAATCATCGCTACAGGTACTTATGACAACACAAATAAACAAATTACCTAC 960
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1051 AATTCTGGAGAAATCATCGCTACAGGTACTTATGACAACACAAATAAACAAATTACCTAC 1110

Qy      961 ACTTTTACAGATTATGTAGATAAATATGAAAATATTAAAGCGCACCTTAAATTAACATCA 1020
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1111 ACTTTTACAGATTATGTAGATAAATATGAAAATATTAAAGCGCACCTTAAATTAACATCA 1170

Qy     1021 TACATTGATAAATCAAAGGTTCCAAATAATAACACTAAGTTAGATGTAGAATATAAGACG 1080
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1171 TACATTGATAAATCAAAGGTTCCAAATAATAACACTAAGTTAGATGTAGAATATAAGACG 1230
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Qy        1081 GCCCTTTCATCAGTAAATAAAACAATTACGGTTGAATATCAAAAACCTAACGAAAATCGG 1140
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1231 GCCCTTTCATCAGTAAATAAAACAATTACGGTTGAATATCAAAAACCTAACGAAAATCGG 1290

Qy        1141 ACTGCTAACCTTCAAAGTATGTTCACAAACATAGATACGAAAAACCATACAGTTGAGCAA 1200
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1291 ACTGCTAACCTTCAAAGTATGTTCACAAACATAGATACGAAAAACCATACAGTTGAGCAA 1350

Qy        1201 ACGATTTATATTAACCCTCTTCGTTATTCAGCCAAAGAAACAAATGTAAATATTTCAGGG 1260
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1351 ACGATTTATATTAACCCTCTTCGTTATTCAGCCAAAGAAACAAATGTAAATATTTCAGGG 1410

Qy        1261 AATGGCGATGAAGGTTCAACAATTATCGACGATAGTACAATCATTAAAGTTTATAAGGTT 1320
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1411 AATGGCGATGAAGGTTCAACAATTATCGACGATAGTACAATCATTAAAGTTTATAAGGTT 1470

Qy        1321 GGAGATAATCAAAATTTACCAGATAGTAACAGAATTTATGATTACAGTGAATATGAAGAT 1380
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1471 GGAGATAATCAAAATTTACCAGATAGTAACAGAATTTATGATTACAGTGAATATGAAGAT 1530

Qy        1381 GTCACAAATGATGATTATGCCCAATTAGGAAATAATAATGACGTGAATATTAATTTTGGT 1440
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1531 GTCACAAATGATGATTATGCCCAATTAGGAAATAATAATGACGTGAATATTAATTTTGGT 1590

Qy        1441 AATATAGATTCACCATATATTATTAAAGTTATTAGTAAATATGACCCTAATAAGGACGAT 1500
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1591 AATATAGATTCACCATATATTATTAAAGTTATTAGTAAATATGACCCTAATAAGGACGAT 1650

Qy        1501 TACACGACGATACAGCAAACTGTGACAATGCAAACGACTATAAATGAGTATACTGGTGAG 1560
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1651 TACACGACGATACAGCAAACTGTGACAATGCAAACGACTATAAATGAGTATACTGGTGAG 1710

Qy        1561 TTTAGAACAGCATCCTATGATAATACAATTGCTTTCTCTACAAGTTCAGGTCAAGGACAA 1620
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1711 TTTAGAACAGCATCCTATGATAATACAATTGCTTTCTCTACAAGTTCAGGTCAAGGACAA 1770

Qy        1621 GGTGACTTGCCTCCTGAAAAAACT 1644
               ||||||||||||||||||||||||
Db        1771 GGTGACTTGCCTCCTGAAAAAACT 1794
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4. Title: US-10-615-383A-16
Sequence: 1 TYTFTDYVD 9
DE Staphylococcus epidermidis ORF amino acid sequence SEQ ID NO:5314.
XX
KW Staphylococcus epidermidis; open reading frame; ORF; bacterial infection;
KW antibacterial; gene therapy.
XX
OS Staphylococcus epidermidis.
XX
PN US6380370-B1.
XX
PD 30-APR-2002.
XX
PF 13-AUG-1998; 98US-00134001.
XX
PR 14-AUG-1997; 97US-0055779P.
PR 08-NOV-1997; 97US-0064964P.
XX
PA (GENO-) GENOME THERAPEUTICS CORP.
XX
PI Doucette-Stamm LA, Bush D;
XX

DR WPI; 2002-381255/41.
DR N-PSDB; ABN93014.
XX
PT Novel isolated nucleic acid encoding a Staphylococcus epidermis
PT polypeptide, useful for diagnosing and treating bacterial infections.
XX
PS Disclosure; SEQ ID NO 5314; 267pp; English.
XX
CC ABN90538 to ABN93374 represent Staphylococcus epidermidis open reading
CC frame (ORF) nucleic acid sequences which encode the amino acid sequences
CC given in ABP35124 to ABP37960. The S. epidermidis sequences have
CC antibacterial activity and can be used in gene therapy. The sequences can
CC also be used in the diagnosis and treatment of bacterial infections,
CC particularly S. epidermidis infections. The sequences can be used to
CC screen for compounds able to interfere with the S. epidermidis life cycle
CC or inhibit S. epidermidis infection. N.B. The sequence data for this
CC patent did not form part of the printed specification, but was obtained
CC in electronic format directly from the USPTO web site
XX
SQ Sequence 930 AA;

Query Match 100.0%; Score 51; DB 1; Length 930;
Best Local Similarity 100.0%; Pred. No. 23;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 TYTFTDYVD 9
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Db        369 TYTFTDYVD 377
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